

BV #12



PCT09

ENTERED

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/889,344B

DATE: 12/04/2002  
 TIME: 14:00:04

Input Set : A:\USSEQLIST3.txt  
 Output Set: N:\CRF4\12032002\I889344B.raw

4 <110> APPLICANT: CHEN, WENFANG  
 5 MEEK, THOMAS D.  
 6 POWELL, DAVID J.  
 7 TEW, DAVID G.  
 10 <120> TITLE OF INVENTION: Method of Site Specific Labeling of Proteins and Uses  
 11 Therefor  
 14 <130> FILE REFERENCE: P50892  
 16 <140> CURRENT APPLICATION NUMBER: 09/889,344B  
 17 <141> CURRENT FILING DATE: 2001-07-16  
 19 <150> PRIOR APPLICATION NUMBER: PCT/US00/01481  
 20 <151> PRIOR FILING DATE: 2000-01-20  
 22 <150> PRIOR APPLICATION NUMBER: US 60/117,327  
 23 <151> PRIOR FILING DATE: 1999-01-22  
 25 <160> NUMBER OF SEQ ID NOS: 16  
 27 <170> SOFTWARE: FastSEQ for Windows Version 3.0  
 29 <210> SEQ ID NO: 1  
 30 <211> LENGTH: 5  
 31 <212> TYPE: PRT  
 32 <213> ORGANISM: Artificial Sequence  
 34 <220> FEATURE:  
 35 <221> NAME/KEY: unsure  
 36 <222> LOCATION: (5)  
 37 <223> OTHER INFORMATION: Where Xaa at position (5) can represent Leucine or Isoleucine  
 39 <220> FEATURE:  
 40 <223> OTHER INFORMATION: Wherein the amino acid sequence is modified by reacting a  
 41 <223> OTHER INFORMATION: transglutaminase with a detectable labeling compound  
 W--> 42 <400> 1  
 W--> 43 Gln Ser Lys Val Xaa  
 44 1 5  
 46 <210> SEQ ID NO: 2  
 47 <211> LENGTH: 207  
 48 <212> TYPE: PRT  
 49 <213> ORGANISM: Artificial Sequence  
 51 <220> FEATURE:  
 52 <221> NAME/KEY: unsure  
 53 <222> LOCATION: (1) (2) (3) (4) (5) (6) (7) (8) (9) (10) (11) (12) (13) (14) (15)  
 54 <222> LOCATION: (16) (17) (18) (19) (20) (21) (22) (23) (24) (25) (26) (27) (28)  
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RAW SEQUENCE LISTING  
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Input Set : A:\USSEQLIST3.txt  
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W--> 61 <222> (110) (111) (112) (113) (114) (115) (116) (117) (118) (119) (120)  
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W--> 64 <222> (143) (144) (145) (146) (147) (148) (149) (150) (151) (152) (153)  
W--> 65 <222> (154) (155) (156) (157) (158) (159) (160) (161) (162) (163) (164)  
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70 <223> OTHER INFORMATION: Where Xaa can represent none or any one of the twenty naturally

71 <223> OTHER INFORMATION: occurring amino acids

W--> 73 <220> FEATURE:

74 <223> OTHER INFORMATION: Wherein the amino acid sequence is modified by reacting a  
75 <223> OTHER INFORMATION: transglutaminase with a detectable labeling compound

W--> 77 <400> 2

W--> 78 Xaa  
79 1 5 10 15

W--> 80 Xaa  
81 20 25 30

W--> 82 Xaa  
83 35 40 45

W--> 84 Xaa  
85 50 55 60

W--> 86 Xaa  
87 65 70 75 80

W--> 88 Xaa  
89 85 90 95

W--> 90 Xaa Xaa Xaa Xaa Gln Ser Lys Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa  
91 100 105 110

W--> 92 Xaa  
93 115 120 125

W--> 94 Xaa  
95 130 135 140

W--> 96 Xaa  
97 145 150 155 160

W--> 98 Xaa  
99 165 170 175

W--> 100 Xaa  
101 180 185 190

W--> 102 Xaa  
103 195 200 205

105 <210> SEQ ID NO: 3

106 <211> LENGTH: 207

107 <212> TYPE: PRT

108 <213> ORGANISM: Artificial Sequence

110 <220> FEATURE:

111 <221> NAME/KEY: unsure

112 <222> LOCATION: (1) (2) (3) (4) (5) (6) (7) (8) (9) (10) (11) (12) (13) (14) (15)

113 <222> LOCATION: (16) (17) (18) (19) (20) (21) (22) (23) (24) (25) (26) (27) (28)

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W--> 114 <222> (29) (30) (31) (32) (33) (34) (35) (36) (37) (38) (39) (40) (41)  
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W--> 116 <222> (55) (56) (57) (58) (59) (60) (61) (62) (62) (64) (65) (66) (67)  
W--> 117 <222> (68) (69) (70) (71) (72) (73) (74) (75) (76) (77) (78) (79) (80)  
W--> 118 <222> (81) (82) (83) (84) (85) (86) (87) (88) (89) (90) (91) (92) (93)  
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W--> 120 <222> (110) (111) (112) (113) (114) (115) (116) (117) (118) (119) (120)  
W--> 121 <222> (121) (122) (123) (124) (125) (126) (127) (128) (129) (130) (131)  
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W--> 123 <222> (143) (144) (145) (146) (147) (148) (149) (150) (151) (152) (153)  
W--> 124 <222> (154) (155) (156) (157) (158) (159) (160) (161) (162) (163) (164)  
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W--> 126 <222> (176) (177) (178) (179) (180) (181) (182) (183) (184) (185) (186)  
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W--> 128 <222> (198) (199) (200) (201) (202) (203) (204) (205) (206) (207)

129 <223> OTHER INFORMATION: Where Xaa can represent none or any one of the twenty naturally

130 <223> OTHER INFORMATION: occurring amino acids

W--> 132 <220> FEATURE:

133 <223> OTHER INFORMATION: Wherein the amino acid sequence is modified by reacting a  
134 <223> OTHER INFORMATION: transglutaminase with a detectable labeling compound

W--> 136 <400> 3

W--> 137	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
138	1											15
W--> 139	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
140		20										30
W--> 141	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
142			35									45
W--> 143	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
144			50									60
W--> 145	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
146			65									80
W--> 147	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
148			85									95
W--> 149	Xaa	Xaa	Xaa	Xaa	Gln	Ser	Lys	Val	Xaa	Xaa	Xaa	Xaa
150					100			105				110
W--> 151	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
152			115					120				125
W--> 153	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
154			130					135				140
W--> 155	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
156			145					150				160
W--> 157	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
158			165					170				175
W--> 159	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
160			180					185				190
W--> 161	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
162			195					200				205
164	<210>	SEQ ID NO:	4									
165	<211>	LENGTH:	10									

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166 <212> TYPE: PRT  
167 <213> ORGANISM: Artificial Sequence  
169 <220> FEATURE:  
170 <223> OTHER INFORMATION: Derivative of a factor XIII substrate  
172 <400> SEQUENCE: 4  
173 Leu Ser Leu Ser Gln Ser Lys Val Leu Gly  
174 1 5 10  
176 <210> SEQ ID NO: 5  
177 <211> LENGTH: 10  
178 <212> TYPE: PRT  
179 <213> ORGANISM: Artificial Sequence  
181 <220> FEATURE:  
182 <223> OTHER INFORMATION: Derivative of a factor XIII substrate  
184 <400> SEQUENCE: 5  
185 Ile Gly Glu Gly Gln Ser Lys Val Leu Gly  
186 1 5 10  
188 <210> SEQ ID NO: 6  
189 <211> LENGTH: 10  
190 <212> TYPE: PRT  
191 <213> ORGANISM: Artificial Sequence  
193 <220> FEATURE:  
194 <223> OTHER INFORMATION: Derivative of a factor XIII substrate  
196 <400> SEQUENCE: 6  
197 Leu Gly Pro Gly Gln Ser Lys Val Ile Gly  
198 1 5 10  
200 <210> SEQ ID NO: 7  
201 <211> LENGTH: 81  
202 <212> TYPE: DNA  
203 <213> ORGANISM: Unknown  
205 <220> FEATURE:  
206 <223> OTHER INFORMATION: Oligonucleotide designed to introduce Q tag  
208 <400> SEQUENCE: 7  
209 tgtacacctcg accatatgag cctgtccctg tcccagtcca aagttctgcc gggtccgagc 60  
210 actatcgaaq aacgcgttaa g 81  
212 <210> SEQ ID NO: 8  
213 <211> LENGTH: 37  
214 <212> TYPE: DNA  
215 <213> ORGANISM: Unknown  
217 <220> FEATURE:  
218 <223> OTHER INFORMATION: Oligonucleotide designed to introduce Q tag  
220 <400> SEQUENCE: 8  
221 tgatgtcaqt caagcttacg cctggtgcc gttgatg 37  
223 <210> SEQ ID NO: 9  
224 <211> LENGTH: 14  
225 <212> TYPE: PRT  
226 <213> ORGANISM: Artificial Sequence  
228 <220> FEATURE:  
229 <223> OTHER INFORMATION: Derivative of a factor XIII substrate  
231 <400> SEQUENCE: 9

## **RAW SEQUENCE LISTING**

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Input Set : A:\USSEQLIST3.txt

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232 Met Ser Leu Ser Leu Ser Gln Ser Lys Val Leu Pro Gly Pro  
233 1 5 10  
235 <210> SEQ ID NO: 10  
236 <211> LENGTH: 37  
237 <212> TYPE: DNA  
238 <213> ORGANISM: Unknown  
240 <220> FEATURE:  
241 <223> OTHER INFORMATION: Oligonucleotide designed to introduce Q tag  
243 <400> SEQUENCE: 10  
244 tgtacctcag accatatgag cactatcgaa gaacgcg 37  
246 <210> SEQ ID NO: 11  
247 <211> LENGTH: 78  
248 <212> TYPE: DNA  
249 <213> ORGANISM: Unknown  
251 <220> FEATURE:  
252 <223> OTHER INFORMATION: Oligonucleotide designed to introduce Q tag  
254 <400> SEQUENCE: 11  
255 ttagtgcagt caagcttacg gaccggcag aactttggac tgggacaggg acagcgctg 60  
256 gtggccgtt atgtaatc 78  
258 <210> SEQ ID NO: 12  
259 <211> LENGTH: 12  
260 <212> TYPE: PRT  
261 <213> ORGANISM: Artificial Sequence  
263 <220> FEATURE:  
264 <223> OTHER INFORMATION: Derivative of E. coli ACP protein  
266 <400> SEQUENCE: 12  
267 Leu Ser Leu Ser Gln Ser Lys Val Leu Pro Gly Pro  
268 1 5 10  
270 <210> SEQ ID NO: 13  
271 <211> LENGTH: 92  
272 <212> TYPE: DNA  
273 <213> ORGANISM: Unknown  
275 <220> FEATURE:  
276 <223> OTHER INFORMATION: Oligonucleotide designed to introduce Q tag into  
277 Streptococcus haemophilus FabH gene 60  
279 <400> SEQUENCE: 13  
280 tatcatatga gcctgtccct gtcccagtcc aaagttctgc cgggtccggg taccctcgag 92  
281 ggatccgctt ttgcaaaaat aagtcaagggtt gc  
283 <210> SEQ ID NO: 14  
284 <211> LENGTH: 53  
285 <212> TYPE: DNA  
286 <213> ORGANISM: Unknown  
288 <220> FEATURE:  
289 <223> OTHER INFORMATION: Oligonucleotide designed to introduce Q tag into  
290 Streptococcus haemophilus FabH gene  
292 <400> SEQUENCE: 14  
293 ctcatatgt agtcactag tggatccta aattgttata atgagcgtgc ccc 53  
295 <210> SEQ ID NO: 15  
296 <211> LENGTH: 364

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/889,344B

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Input Set : A:\USSEQLIST3.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 5  
Seq#:2; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22  
Seq#:2; Xaa Pos. 23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41  
Seq#:2; Xaa Pos. 42,43,44,45,46,47,48,49,50,51,52,53,54,55,56,57,58,59,60  
Seq#:2; Xaa Pos. 61,62,63,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79  
Seq#:2; Xaa Pos. 80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98  
Seq#:2; Xaa Pos. 99,100,101,106,107,108,109,110,111,112,113,114,115,116,117  
Seq#:2; Xaa Pos. 118,119,120,121,122,123,124,125,126,127,128,129,130,131  
Seq#:2; Xaa Pos. 132,133,134,135,136,137,138,139,140,141,142,143,144,145  
Seq#:2; Xaa Pos. 146,147,148,149,150,151,152,153,154,155,156,157,158,159  
Seq#:2; Xaa Pos. 160,161,162,163,164,165,166,167,168,169,170,171,172,173  
Seq#:2; Xaa Pos. 174,175,176,177,178,179,180,181,182,183,184,185,186,187  
Seq#:2; Xaa Pos. 188,189,190,191,192,193,194,195,196,197,198,199,200,201  
Seq#:2; Xaa Pos. 202,203,204,205,206,207  
Seq#:3; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22  
Seq#:3; Xaa Pos. 23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41  
Seq#:3; Xaa Pos. 42,43,44,45,46,47,48,49,50,51,52,53,54,55,56,57,58,59,60  
Seq#:3; Xaa Pos. 61,62,63,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79  
Seq#:3; Xaa Pos. 80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98  
Seq#:3; Xaa Pos. 99,100,101,106,107,108,109,110,111,112,113,114,115,116,117  
Seq#:3; Xaa Pos. 118,119,120,121,122,123,124,125,126,127,128,129,130,131  
Seq#:3; Xaa Pos. 132,133,134,135,136,137,138,139,140,141,142,143,144,145  
Seq#:3; Xaa Pos. 146,147,148,149,150,151,152,153,154,155,156,157,158,159  
Seq#:3; Xaa Pos. 160,161,162,163,164,165,166,167,168,169,170,171,172,173  
Seq#:3; Xaa Pos. 174,175,176,177,178,179,180,181,182,183,184,185,186,187  
Seq#:3; Xaa Pos. 188,189,190,191,192,193,194,195,196,197,198,199,200,201  
Seq#:3; Xaa Pos. 202,203,204,205,206,207

VERIFICATION SUMMARY  
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Input Set : A:\USSEQLIST3.txt  
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L:42 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1  
L:42 M:283 W: Missing Blank Line separator, <400> field identifier  
L:43 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:55 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:56 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:57 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:58 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:59 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:60 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:61 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:62 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:63 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:64 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:65 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:66 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
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L:68 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:69 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:77 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2  
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:16  
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:32  
L:84 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:48  
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:64  
L:88 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:80  
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:96  
L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:112  
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:128  
L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:144  
L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:160  
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:176  
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:192  
L:114 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:115 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:116 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
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L:121 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:122 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:123 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
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L:125 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:126 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:127 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:128 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3  
L:136 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3

**VERIFICATION SUMMARY**

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L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:16  
L:141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:32  
L:143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:48  
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:64  
L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:80  
L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:96  
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:112  
L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:128  
L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:144  
L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:160  
L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:176  
L:161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:192